```
In [62]:
          import numpy as np
          import sklearn as skl
          from sklearn import datasets
          from sklearn.model_selection import train_test_split
          from sklearn.preprocessing import MinMaxScaler
          from sklearn.metrics import classification_report
          import pandas as pd
          from numpy import linalg
          from scipy.special import expit
 In [3]:
          data=datasets.load_breast_cancer()
          df = pd.DataFrame(data.data, columns=data.feature names)
          df['target'] = data.target
In [58]:
          def SGD(data,validate,epsilon=0.1,target=None,batch_size=None,alpha=0.005,max_it
              if batch size is None:
                  batch_size=int(0.1*data.shape[0])
                  #sets batch size to 5% of population if none entered
              w=np.random.normal(0,1,data.shape[1])
              #generates initial weights and biases for model from standard normal
              for i in np.arange(max iter):
                  sample=data.sample(n=batch_size)
                  #samples data from population
                  x=np.insert(sample.iloc[:,:-1].values,0,np.ones([batch_size,]),axis=1)
                  #adds column of ones for w_0 of weights and bias vector
                  t=sample.iloc[:,-1].values
                  #vector of targets
                  y=expit(x.dot(w))
                  #compute the y_n for each x_n
                  g=np.array([diff*phi for diff,phi in zip(y-t,x)]).sum(axis=0)
                  #compute the gradient
                  w=w-alpha*g
                  #compute new weight and bias vector
                  if i%1000==0:
                      x_v=np.insert(validate.iloc[:,:-1].values,0,np.ones([validate.shape[
                      y_v=expit(x_v.dot(w))
                      t_v=validate.iloc[:,-1].values
                      error=np.sqrt(((y_v - t_v) ** 2).mean())
                      if error<epsilon:</pre>
                          break
              return w, error, i #return w vector, RMSE, and iterations
In [31]:
          scaler=MinMaxScaler()
          scaled_data = scaler.fit_transform(df)
          scaled_df = pd.DataFrame(scaled_data, columns=df.columns)
In [76]:
          scaled_df.head()
```

| Out[76]: | | mean radius | mean texture | mean perimeter | mea are | | mean compactness | mean concavity | mean concave points | syn | |
|---------------------|---|--|-----------------|-------------------|--------------|----------------------|---------------------|-------------------|---------------------------|-----|--|
| | 0 | 0.521037 | 0.022658 | 0.545989 | 0.36373 | 3 0.593753 | 0.792037 | 0.703140 | 0.731113 | 0.6 | |
| | 1 | 0.643144 | 0.272574 | 0.615783 | 0.50159 | 0.289880 | 0.181768 | 0.203608 | 0.348757 | 0. | |
| | 2 | 0.601496 | 0.390260 | 0.595743 | 0.44941 | 7 0.514309 | 0.431017 | 0.462512 | 0.635686 | 3.0 | |
| | 3 | 0.210090 | 0.360839 | 0.233501 | 0.10290 | 6 0.811321 | 0.811361 | 0.565604 | 0.522863 | 0. | |
| | 4 | 0.629893 | 0.156578 | 0.630986 | 0.48929 | 0.430351 | 0.347893 | 0.463918 | 0.518390 | 0.: | |
| 5 rows × 31 columns | | | | | | | | | | | |
| In [38]: | train, validate, test = np.split(scaled_df.sample(frac=1), [int(.6*len(scaled_df.sample)) | | | | | | | | | | |
| In [61]: | t | <pre>test_features=np.insert(test.iloc[:,:-1].values,0,np.ones([test.shape[0],]),axis test_target=test.iloc[:,-1].values predicted_target=expit(test_features.dot(SGD(train,validate)[0]))</pre> | | | | | | | | | |
| In [75]: | p | <pre>print(classification_report(test_target,np.round(predicted_target)))</pre> | | | | | | | | | |
| | | | preci | sion r | ecall · | f1-score s | upport | | | | |
| | | | | 0.93 0.91 | 0.88 0.95 | 0.91 0.93 | 49 65 | | | | |
| | we | accurad macro av ighted av | /g | 0.92 0.92 | 0.92 0.92 | 0.92 0.92 0.92 | 114 114 114 | | | | |